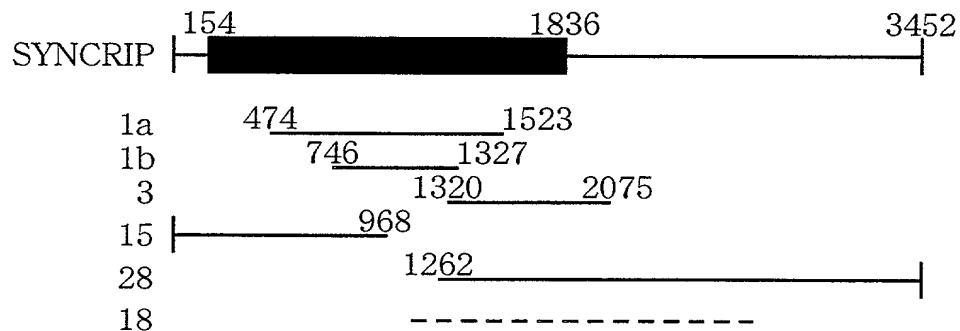


FIG.1

A



B

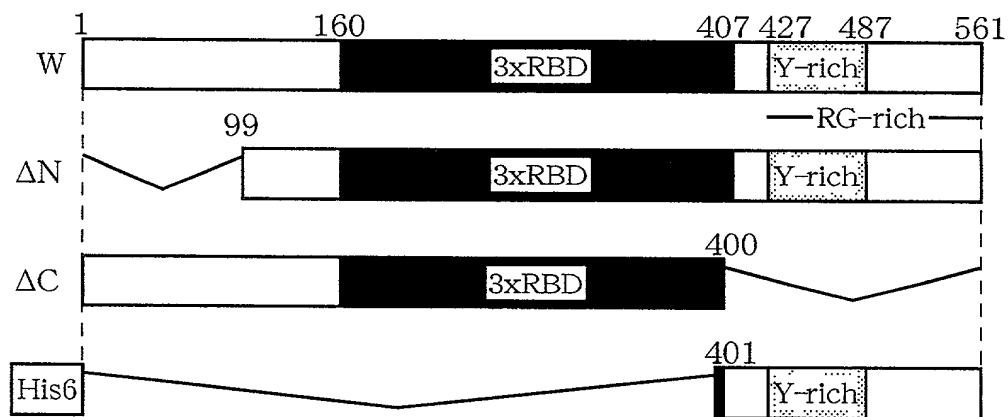


FIG.2

SYNCRIP	1: ---MATEHVNGNGTTEFMDTTSAVIRSENFQTLDDAGLPQKVAEKLDSEIYVAGLVANSD	56
hnRNPR	1: MANQVNGNAVQLKEESEPMD-TSSVTHTEHYKTLIEAGLPQKVAERLDEIFQTLGLVAVVD	59
SYNCRIP	57: LDERATEALKEFNEDGALAVLQOFKDSHLSHVQNKSAFLGGVMKTYRQREKOGTKVADSS	116
hnRNPR	60: LDERAIDALREFNEEGALSVLQOFKESDLSHVQNKSAFLCGVMKTYRQREKOGSKVQEST	119
SYNCRIP	117: KGPDEAKIKALLERTGYTLDTVTGQRKYGGPPEDSVYSGQQPSVGTEIFVGKIERDLTFED	176
hnRNPR	120: KGPDEAKIKALLERTGYTLDTVTGQRKYGGPPEDSVYSGVOPGIGTEVEVGKIERDLTFED	179
SYNCRIP	177: ELVPLFERAGPIWDLRLMMDPLTGLNRGYAFVTFCTKEAAQBAVKLYNMHEIRSGKHIGV	236
hnRNPR	180: ELVPLFERAGPIWDLRLMMDPLSGQNRGYAFITFCGKEAAQBAVKLCDSYERPGKHIGV	239
SYNCRIP	237: CISVANNRLEFVSGTEKSKTKKQILEEFSKVTEGLTDVILYHQPDDKKKNRGFCFLEYEDH	296
hnRNPR	240: CISVANNRLEFVSGTEKSKTKRENILEEFSKVTEGLVDVILYHQPDDKKKNRGFCFLEYEDH	299
clone-18 (m-hnRNPR)	1: -----DH	2
SYNCRIP	297: KTAQAARRRLMSCKVKVWGNVGVTEWADPIEDDPVMAKVKVLEVRNLANTVTETELK	356
hnRNPR	300: KSAAQAARRRLMSCKVKVWGNVGVTEWADPVVEEDPVMAKVKVLEVRNLATTVTETELK	359
clone-18 (m-hnRNPR)	3: KSAAQAARRRLMSCKVKVWGNVGVTEWADPVVEEDPVMAKVKVLEVRNLATTVTETELK	62
SYNCRIP	357: SESEFGKLERVKIKDYAFVHFEEDRGAAVKAMDEHNGKEIEGEEIEIVLAKPPDKKKER	416
hnRNPR	360: SESEFGKLERVKIKDYAFVHFEEDRGAAVKAMDEHNGKEIEGEEIEIVLAKPPDKKKER	419
clone-18 (m-hnRNPR)	63: SESEFGKLERVKIKDYAFVHFEEDRGAAVKAMDEHNGKEIEGEEIEIVLAKPPDKKKER	122
SYNCRIP	417: KAQRQAANKQMYDDYYYYGPPHPPPTRGRGR-GGRGGYGYPEDYGYEDYY-DYGYDY	474
hnRNPR	420: QAARQASRSTAYEDYYHPPPRMPPPIRGRGRGGGRGGYGYPEDYGYEDYYDDYGYDY	479
clone-18 (m-hnRNPR)	123: QAARQASRSTAYEDYYHPPPRMPPPMRGRGR-GGRGGYGYPEDYGYEDYYDDYGYDY	181
SYNCRIP	475: HNYRGGYEDPYGYED-FQVGARGRGGRGARGA-APSRGRGAAPPGRGAGYSQRGGP-GS	531
hnRNPR	480: HDYRGGYEDPYGYDDGYAVRGGR-GGRGGRGAPPFRGRGAPPFRGAGYSQRGAPLGP	538
clone-18 (m-hnRNPR)	182: HDYRGGYEDPYGYDDGYAVRGGR-GGRGGRGAPPFRGRGAPPFRGAGYSQRGAPLGP	240
SYNCRIP	532: ARGVRGARGG-AQQQRGRGGKGV-EA-GPDLLQ-----	561
hnRNPR	539: PRGSRGGRGPPAQQQRGRGSRGSRGNRGGNVGGRKADGYNQPDSEKRRDTNNQONWGSQP	598
clone-18 (m-hnRNPR)	241: PRGSRGGRGPPAQQQRGRGSRGARGNRGGNVGGRKADGYNQPDSEKRRDTNNQONWGSQP	300
hnRNPR	562: -----	
clone-18 (m-hnRNPR)	599: IAQQPLQGGSDYSGNYGYNNDNQEFYQDTYGQQWK	633
	301: IAQQPLQGGSDYSGNYGYNNDNQEFYQDTYGQQWK	335

FIG.3

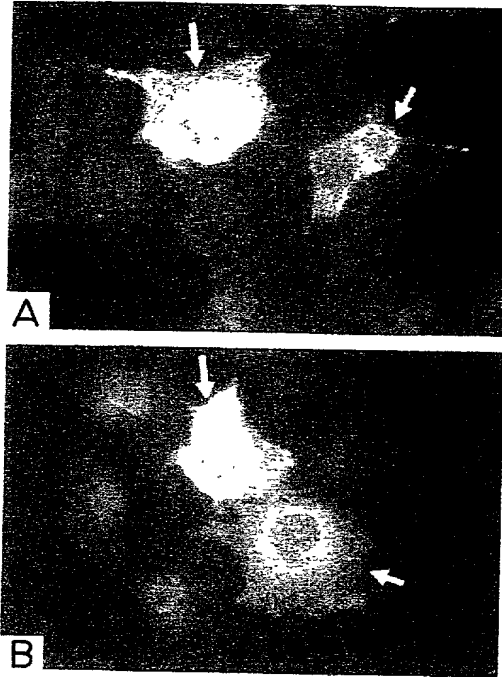


FIG.4

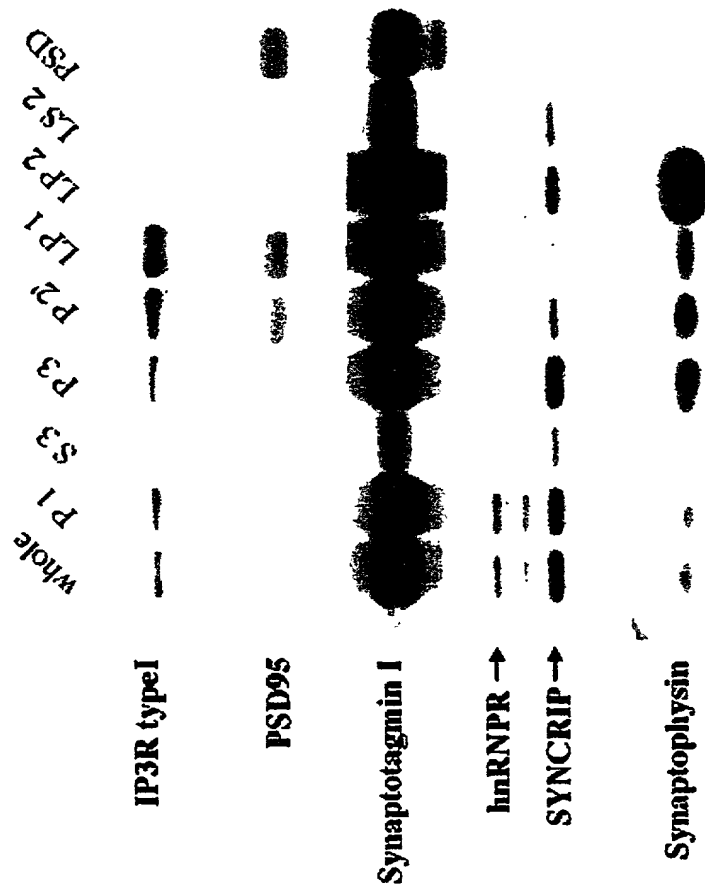


FIG.5

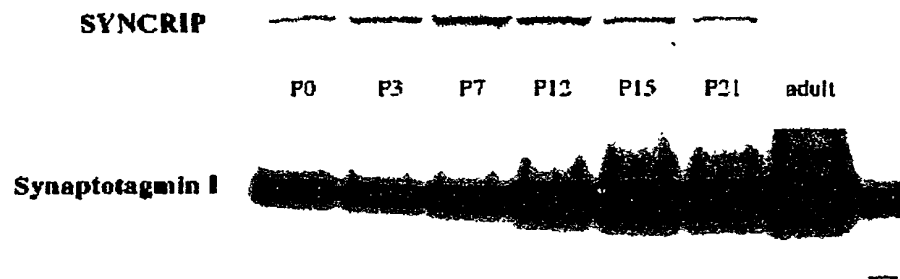


FIG.6

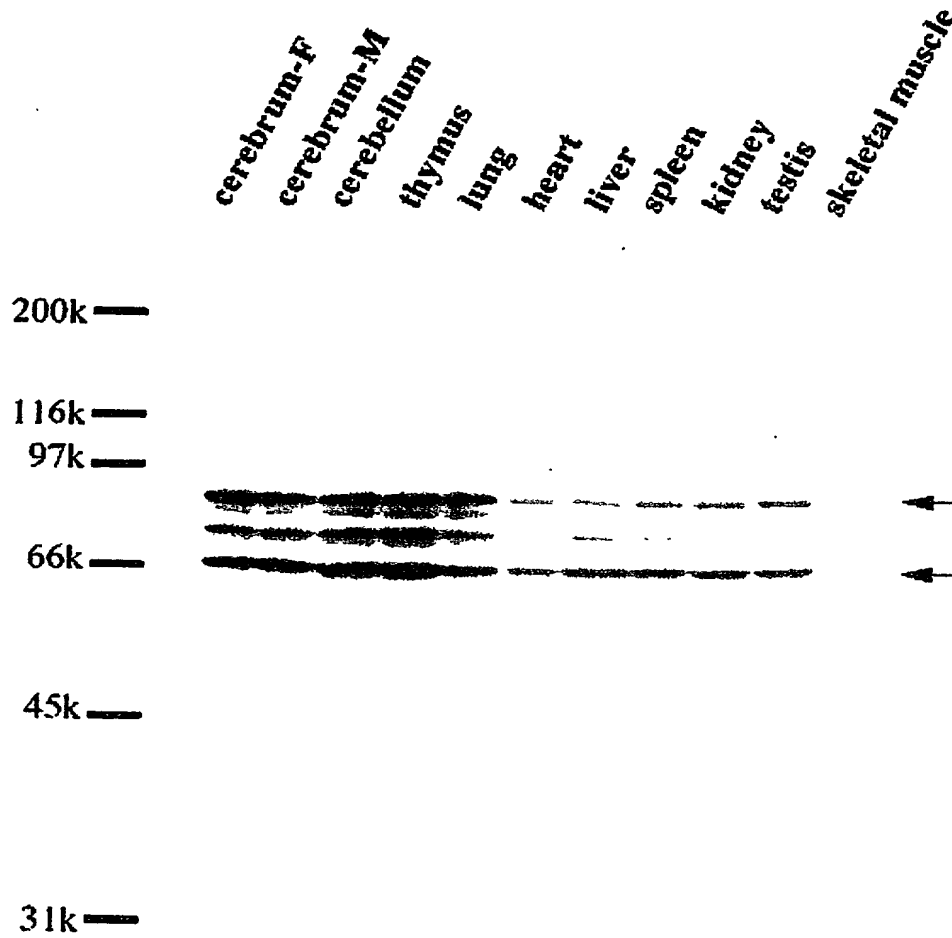


FIG. 7

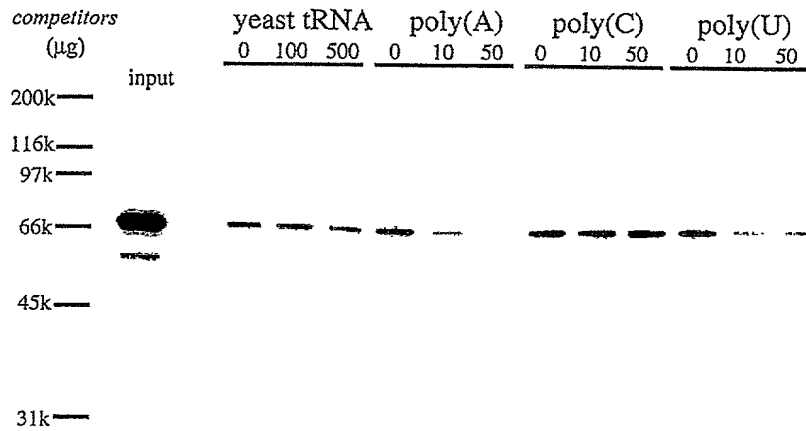


FIG.8

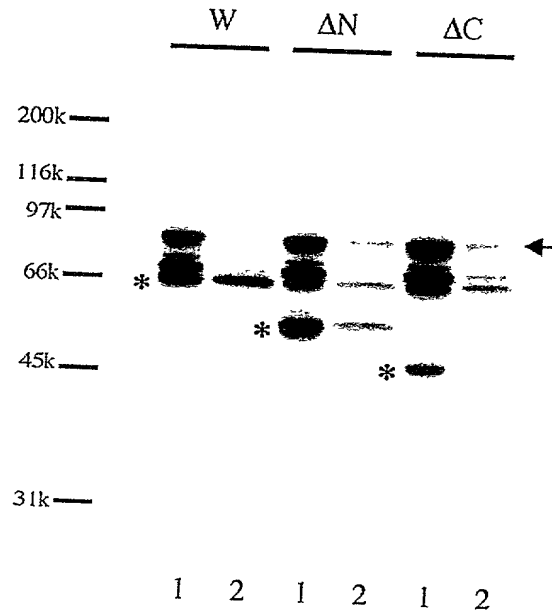


FIG.9

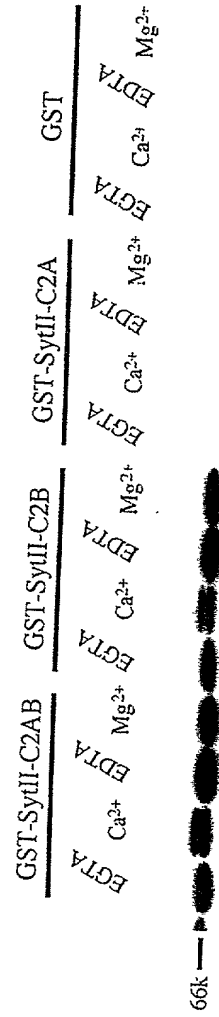


FIG.10

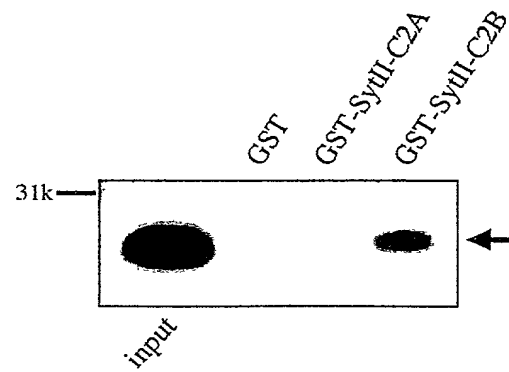


FIG.11

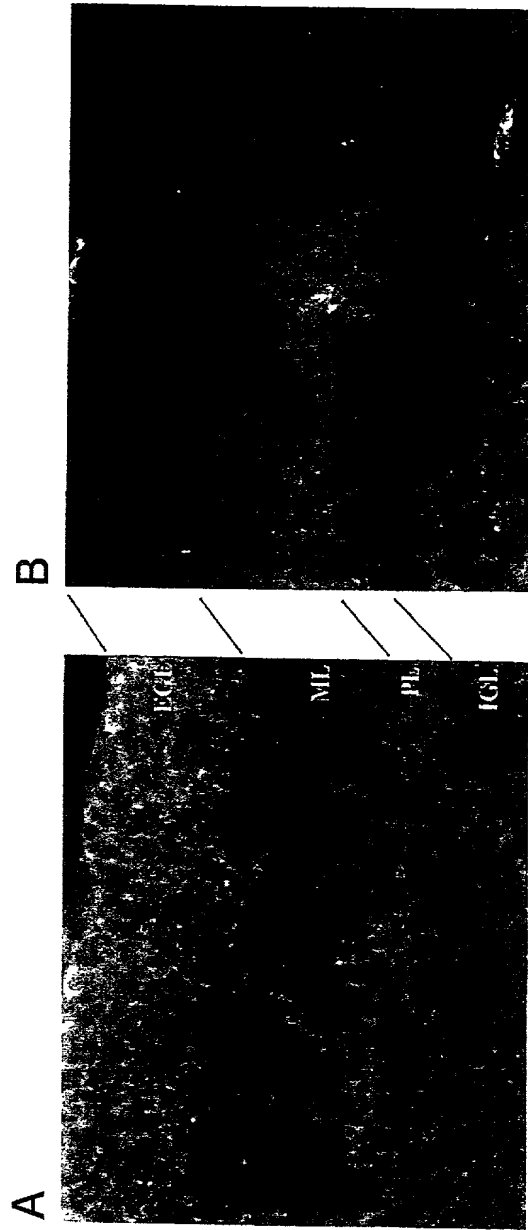


FIG.12

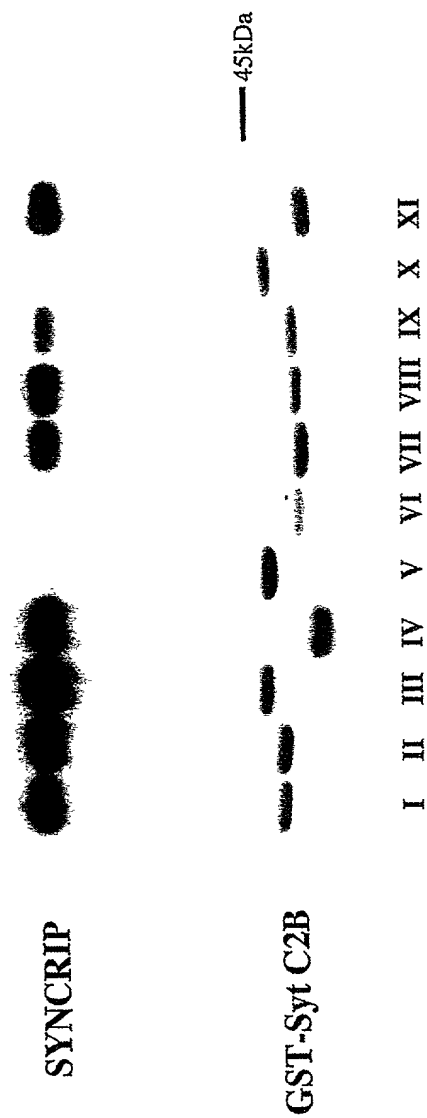


FIG.13

